

2/11 #2



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RAW SEQUENCE LISTING DATE: 02/19/2002
 PATENT APPLICATION: US/10/057,113 TIME: 15:10:40

Input Set : N:\Crf3\RULE60\10057113.txt
 Output Set: N:\CRF3\02192002\J057113.raw

SEQUENCE LISTING

3 (i) GENERAL INFORMATION:

5 (i) APPLICANT: THE UNITED STATES OF AMERICA, AS REPRESENTED
 6 BY THE SECRETARY, DEPARTMENT OF HEALTH AND HUMAN SERVICES IN
 7 CARE OF NATIONAL INSTITUTES OF HEALTH Office of Technology
 8 Transfer Suite 325, 6011 Executive Boulevard, Rockville, MD
 9 20852

11 (ii) TITLE OF INVENTION: GLYCOPROTEIN HORMONE SUPERAGONISTS

13 (iii) NUMBER OF SEQUENCES: 37

15 (iv) CORRESPONDENCE ADDRESS:

16 (A) ADDRESSEE: NEEDLE & ROSENBERG, P.C.
 17 (B) STREET: Suite 1200, 127 Peachtree Street
 18 (C) CITY: Atlanta
 19 (D) STATE: Georgia
 20 (E) COUNTRY: USA
 21 (F) ZIP: 30303

23 (v) COMPUTER READABLE FORM:

24 (A) MEDIUM TYPE: Floppy disk
 25 (B) COMPUTER: IBM PC compatible
 26 (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 27 (D) SOFTWARE: PatentIn Release #1.0, Version #1.30

29 (vi) CURRENT APPLICATION DATA:

C--> 30 (A) APPLICATION NUMBER: US/10/057,113
 C--> 31 (B) FILING DATE: 25-Jan-2002
 32 (C) CLASSIFICATION:

34 (vii) PRIOR APPLICATION DATA:

35 (A) APPLICATION NUMBER: PCT/US96/06483
 36 (B) FILING DATE:

40 (viii) ATTORNEY/AGENT INFORMATION:

41 (A) NAME: Perryman, David G.
 42 (B) REGISTRATION NUMBER: 33,438
 43 (C) REFERENCE/DOCKET NUMBER: 14014.0185/P

45 (ix) TELECOMMUNICATION INFORMATION:

46 (A) TELEPHONE: 404/688-0770
 47 (B) TELEFAX: 404/688-9880

50 (2) INFORMATION FOR SEQ ID NO: 1:

52 (i) SEQUENCE CHARACTERISTICS:

53 (A) LENGTH: 22 base pairs
 54 (B) TYPE: nucleic acid
 55 (C) STRANDEDNESS: single
 56 (D) TOPOLOGY: linear

58 (ii) MOLECULE TYPE: DNA (genomic)

63 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

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65 CCTGATAGAT TGCCCAGAAT GC 22
67 (2) INFORMATION FOR SEQ ID NO: 2:
69     (i) SEQUENCE CHARACTERISTICS:
70         (A) LENGTH: 24 base pairs
71         (B) TYPE: nucleic acid
72         (C) STRANDEDNESS: single
73         (D) TOPOLOGY: linear
75     (ii) MOLECULE TYPE: DNA (genomic)
80     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
82 GTGATAATAA CAAGTACTGC AGTG 24
84 (2) INFORMATION FOR SEQ ID NO: 3:
86     (i) SEQUENCE CHARACTERISTICS:
87         (A) LENGTH: 12 amino acids
88         (B) TYPE: amino acid
89         (C) STRANDEDNESS: Not Relevant
90         (D) TOPOLOGY: linear
92     (ii) MOLECULE TYPE: peptide
97     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
99     Cys Thr Leu Gln Glu Asn Pro Phe Phe Ser Gln Pro
100     1           5           10
102 (2) INFORMATION FOR SEQ ID NO: 4:
104     (i) SEQUENCE CHARACTERISTICS:
105         (A) LENGTH: 12 amino acids
106         (B) TYPE: amino acid
107         (C) STRANDEDNESS: Not Relevant
108         (D) TOPOLOGY: linear
110     (ii) MOLECULE TYPE: peptide
115     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
117     Cys Thr Leu Gln Glu Asn Pro Phe Phe Ser Gln Pro
118     1           5           10
120 (2) INFORMATION FOR SEQ ID NO: 5:
122     (i) SEQUENCE CHARACTERISTICS:
123         (A) LENGTH: 12 amino acids
124         (B) TYPE: amino acid
125         (C) STRANDEDNESS: Not Relevant
126         (D) TOPOLOGY: linear
128     (ii) MOLECULE TYPE: peptide
133     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
135     Cys Thr Leu Gln Glu Asn Pro Phe Phe Ser Gln Pro
136     1           5           10
138 (2) INFORMATION FOR SEQ ID NO: 6:
140     (i) SEQUENCE CHARACTERISTICS:
141         (A) LENGTH: 12 amino acids
142         (B) TYPE: amino acid
143         (C) STRANDEDNESS: Not Relevant
144         (D) TOPOLOGY: linear
146     (ii) MOLECULE TYPE: peptide
151     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
153     Cys Gln Leu His Glu Asn Pro Phe Phe Ser Gln Pro

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154      1          5          10
156 (2) INFORMATION FOR SEQ ID NO: 7:
158     (i) SEQUENCE CHARACTERISTICS:
159         (A) LENGTH: 12 amino acids
160         (B) TYPE: amino acid
161         (C) STRANDEDNESS: Not Relevant
162         (D) TOPOLOGY: linear
164     (ii) MOLECULE TYPE: peptide
169     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
171     Cys Lys Pro Arg Glu Asn Gln Phe Phe Ser Lys Pro
172      1          5          10
174 (2) INFORMATION FOR SEQ ID NO: 8:
176     (i) SEQUENCE CHARACTERISTICS:
177         (A) LENGTH: 12 amino acids
178         (B) TYPE: amino acid
179         (C) STRANDEDNESS: Not Relevant
180         (D) TOPOLOGY: linear
182     (ii) MOLECULE TYPE: peptide
187     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
189     Cys Lys Pro Arg Glu Asn Lys Phe Phe Ser Lys Pro
190      1          5          10
192 (2) INFORMATION FOR SEQ ID NO: 9:
194     (i) SEQUENCE CHARACTERISTICS:
195         (A) LENGTH: 12 amino acids
196         (B) TYPE: amino acid
197         (C) STRANDEDNESS: Not Relevant
198         (D) TOPOLOGY: linear
200     (ii) MOLECULE TYPE: peptide
205     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
207     Cys Lys Leu Lys Glu Asn Lys Tyr Phe Ser Arg Leu
208      1          5          10
210 (2) INFORMATION FOR SEQ ID NO: 10:
212     (i) SEQUENCE CHARACTERISTICS:
213         (A) LENGTH: 12 amino acids
214         (B) TYPE: amino acid
215         (C) STRANDEDNESS: Not Relevant
216         (D) TOPOLOGY: linear
218     (ii) MOLECULE TYPE: peptide
223     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
225     Cys Lys Leu Lys Glu Asn Lys Tyr Phe Ser Lys Pro
226      1          5          10
228 (2) INFORMATION FOR SEQ ID NO: 11:
230     (i) SEQUENCE CHARACTERISTICS:
231         (A) LENGTH: 12 amino acids
232         (B) TYPE: amino acid
233         (C) STRANDEDNESS: Not Relevant
234         (D) TOPOLOGY: linear
236     (ii) MOLECULE TYPE: peptide
241     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:

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243      Cys Lys Leu Lys Glu Asn Lys Tyr Phe Ser Lys Pro
244      1              5              10
246 (2) INFORMATION FOR SEQ ID NO: 12:
248      (i) SEQUENCE CHARACTERISTICS:
249          (A) LENGTH: 12 amino acids
250          (B) TYPE: amino acid
251          (C) STRANDEDNESS: Not Relevant
252          (D) TOPOLOGY: linear
254      (ii) MOLECULE TYPE: peptide
259      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:
261      Cys Lys Leu Arg Glu Asn Lys Tyr Phe Phe Lys Leu
262      1              5              10
264 (2) INFORMATION FOR SEQ ID NO: 13:
266      (i) SEQUENCE CHARACTERISTICS:
267          (A) LENGTH: 12 amino acids
268          (B) TYPE: amino acid
269          (C) STRANDEDNESS: Not Relevant
270          (D) TOPOLOGY: linear
272      (ii) MOLECULE TYPE: peptide
277      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:
279      Cys Lys Leu Lys Glu Asn Lys Tyr Phe Ser Lys Leu
280      1              5              10
282 (2) INFORMATION FOR SEQ ID NO: 14:
284      (i) SEQUENCE CHARACTERISTICS:
285          (A) LENGTH: 12 amino acids
286          (B) TYPE: amino acid
287          (C) STRANDEDNESS: Not Relevant
288          (D) TOPOLOGY: linear
290      (ii) MOLECULE TYPE: peptide
295      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:
297      Cys Lys Leu Lys Glu Asn Lys Tyr Phe Ser Lys Leu
298      1              5              10
300 (2) INFORMATION FOR SEQ ID NO: 15:
302      (i) SEQUENCE CHARACTERISTICS:
303          (A) LENGTH: 12 amino acids
304          (B) TYPE: amino acid
305          (C) STRANDEDNESS: Not Relevant
306          (D) TOPOLOGY: linear
308      (ii) MOLECULE TYPE: peptide
313      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:
315      Cys Lys Leu Lys Glu Asn Lys Tyr Phe Ser Lys Leu
316      1              5              10
318 (2) INFORMATION FOR SEQ ID NO: 16:
320      (i) SEQUENCE CHARACTERISTICS:
321          (A) LENGTH: 12 amino acids
322          (B) TYPE: amino acid
323          (C) STRANDEDNESS: Not Relevant
324          (D) TOPOLOGY: linear
326      (ii) MOLECULE TYPE: peptide

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331 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:
333 Cys Lys Leu Lys Glu Asn Lys Tyr Phe Ser Lys Leu
334 1 5 10

336 (2) INFORMATION FOR SEQ ID NO: 17:
338 (i) SEQUENCE CHARACTERISTICS:
339 (A) LENGTH: 12 amino acids
340 (B) TYPE: amino acid
341 (C) STRANDEDNESS: Not Relevant
342 (D) TOPOLOGY: linear
344 (ii) MOLECULE TYPE: peptide

349 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:
351 Cys Lys Leu Lys Gln Asn Lys Tyr Phe Ser Lys Leu
352 1 5 10

354 (2) INFORMATION FOR SEQ ID NO: 18:
356 (i) SEQUENCE CHARACTERISTICS:
357 (A) LENGTH: 12 amino acids
358 (B) TYPE: amino acid
359 (C) STRANDEDNESS: Not Relevant
360 (D) TOPOLOGY: linear
362 (ii) MOLECULE TYPE: peptide

367 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:
369 Cys Lys Leu Gly Glu Asn Arg Phe Phe Ser Lys Pro
370 1 5 10

372 (2) INFORMATION FOR SEQ ID NO: 19:
374 (i) SEQUENCE CHARACTERISTICS:
375 (A) LENGTH: 12 amino acids
376 (B) TYPE: amino acid
377 (C) STRANDEDNESS: Not Relevant
378 (D) TOPOLOGY: linear
380 (ii) MOLECULE TYPE: peptide

385 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:
387 Cys Lys Leu Gly Glu Asn Arg Phe Phe Ser Lys Pro
388 1 5 10

390 (2) INFORMATION FOR SEQ ID NO: 20:
392 (i) SEQUENCE CHARACTERISTICS:
393 (A) LENGTH: 12 amino acids
394 (B) TYPE: amino acid
395 (C) STRANDEDNESS: Not Relevant
396 (D) TOPOLOGY: linear
398 (ii) MOLECULE TYPE: peptide

403 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:
405 Cys Lys Leu Gly Glu Asn Arg Phe Phe Ser Lys Pro
406 1 5 10

408 (2) INFORMATION FOR SEQ ID NO: 21:
410 (i) SEQUENCE CHARACTERISTICS:
411 (A) LENGTH: 12 amino acids
412 (B) TYPE: amino acid
413 (C) STRANDEDNESS: Not Relevant
414 (D) TOPOLOGY: linear

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/057,113

DATE: 02/19/2002

TIME: 15:10:41

Input Set : N:\Crf3\RULE60\10057113.txt

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L:30 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]

L:31 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]